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1600

RAW SEQUENCE LISTING

DATE: 07/09/2003

PATENT APPLICATION: US/09/830,111D

TIME: 09:22:53

Input Set : A:\21581-265.seq.ST25.txt
Output Set: N:\CRF4\07092003\I830111D.raw

3 <110> APPLICANT: Kaneka Corporation
4 Matsuda, Hideyuki
5 Kawamukai, Makota
6 Yajima, Kazuyoshi
7 Ikenaka, Yasuhiro
8 Hasegawa, Junzo
9 Takahashi, Satomi
11 <120> TITLE OF INVENTION: Process For Producing Coenzyme Q10
13 <130> FILE REFERENCE: 21581-00265-US
15 <140> CURRENT APPLICATION NUMBER: 09/830,111D
16 <141> CURRENT FILING DATE: 2001-07-23
18 <160> NUMBER OF SEQ ID NOS: 2
20 <170> SOFTWARE: PatentIn version 3.2
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 1653
24 <212> TYPE: DNA
25 <213> ORGANISM: Saioella complicata

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 100 Arg Leu Arg Cys Thr Pro Thr Ser Arg Pro Ser Ser Trp Ala Ala
 101 35 40 45
 104 Ala Val Ser Ser Ala Ser Arg Leu Val Glu Pro Asp Pro Asn Gln Pro
 105 50 55 60
 108 Leu Ile Asn Pro Leu Asn Leu Val Gly Pro Glu Met Ser Asn Leu Thr
 109 65 70 75 80
 112 Ser Asn Ile Arg Ser Leu Leu Gly Ser Gly His Pro Ser Leu Asp Thr
 113 85 90 95
 116 Val Ala Lys Tyr Tyr Val Gln Ser Glu Gly Lys His Ile Arg Pro Leu
 117 100 105 110
 120 Met Val Leu Leu Met Ala Gln Ala Thr Glu Val Ala Pro Lys Val Gln
 121 115 120 125
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 125 130 135 140
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 129 145 150 155 160
 132 Gly Pro Leu Thr Lys Asp Gly Glu Ile Glu Gly Gln Thr Ser Asn Ile
 133 165 170 175
 136 Leu Ala Ser Gln Arg Arg Leu Ala Glu Ile Thr Glu Met Ile His Ala
 137 180 185 190
 140 Ala Ser Leu Leu His Asp Asp Val Ile Asp Ala Ser Glu Thr Arg Arg
 141 195 200 205
 144 Asn Ala Pro Ser Gly Asn Gln Ala Phe Gly Asn Lys Met Ala Ile Leu
 145 210 215 220
 148 Ala Gly Asp Phe Leu Leu Gly Arg Ala Ser Val Ala Leu Ala Arg Leu
 149 225 230 235 240
 152 Arg Asn Pro Glu Val Ile Glu Leu Leu Ala Thr Val Ile Ala Asn Leu
 153 245 250 255
 156 Val Glu Gly Glu Phe Met Gln Leu Lys Asn Thr Val Asp Asp Ala Ile
 157 260 265 270
 160 Glu Ala Thr Ala Thr Gln Glu Thr Phe Asp Tyr Tyr Leu Gln Lys Thr
 161 275 280 285
 164 Tyr Leu Lys Thr Ala Ser Leu Ile Ala Lys Ser Cys Arg Ala Ser Ala
 165 290 295 300
 168 Leu Leu Gly Gly Ala Thr Pro Glu Val Ala Asp Ala Ala Tyr Ala Tyr
 169 305 310 315 320
 172 Gly Arg Asn Leu Gly Leu Ala Phe Gln Ile Val Asp Asp Met Leu Asp

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177 340 345 350
180 Gln Leu Gly Leu Ala Thr Ala Pro Ala Leu Phe Ala Trp Lys His His
181 355 360 365
184 Ala Glu Leu Gly Pro Met Ile Lys Arg Lys Phe Ser Asp Pro Gly Asp
185 370 375 380
188 Val Glu Arg Ala Arg Glu Leu Val Glu Lys Ser Asp Gly Leu Glu Lys
189 385 390 395 400
192 Thr Arg Ala Leu Ala Glu Glu Tyr Ala Gln Lys Ala Leu Asp Ala Ile
193 405 410 415
196 Arg Thr Phe Pro Glu Ser Pro Ala Arg Lys Ala Leu Glu Gln Leu Thr
197 420 425 430
200 Asp Lys Val Leu Thr Arg Ser Arg
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VERIFICATION SUMMARY

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